

A) has six disulphide bonds that form two trefoil domains, where the 12 cysteines that form the six disulphide bonds are in the configuration 1-5, 2-4, 3-6, 7-11, 8-10 and 9-12, and

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B) is encoded by a nucleic acid sequence that is at least 60 % homologous to a nucleic acid sequence that encodes SEQ ID NO:1 and that hybridizes under high stringency conditions to the nucleic acid sequence that encodes SEQ ID NO:1.

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Please add the following claims:

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42. (New) The polypeptide of claim 27, wherein the polypeptide has an amino acid sequence that is a homologue of SEQ ID NO:1.

43. (New) The polypeptide of claim 42, wherein the homologue amino acid sequence contains 39 amino acids in the first trefoil domain.

44. (New) The polypeptide of claim 43, wherein the 39 amino acids in the first trefoil domain of the homologue amino acid sequence are identical to amino acids 8-46 of SEQ ID NO:1 except for two amino acid substitutions.

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45. (New) The polypeptide of claim 44, wherein the 39 amino acids in the first trefoil domain of the homologue amino acid sequence are identical to amino acids 8-46 of SEQ ID NO:1.

46. (New) The polypeptide of claim 42, wherein the homologue amino acid sequence contains 38 amino acids in the second trefoil domain.

47. (New) The polypeptide of claim 46, wherein the 38 amino acids in the second trefoil domain of the homologue amino acid sequence are identical to amino acids 58-95 of SEQ ID NO:1 except for two amino acid substitutions.

48. (New) The polypeptide of claim 47, wherein the 38 amino acids in the second trefoil domain of the homologue amino acid sequence are identical to amino acids 58-95 of SEQ ID NO:1.

49. (New) The polypeptide of claim 43, wherein the homologue amino acid sequence contains 38 amino acids in the second trefoil domain and the 38 amino acids in the second trefoil domain of the homologue amino acid sequence are identical to amino acids 58-95 of SEQ ID NO:1 except for two amino acid substitutions.

50. (New) The polypeptide of claim 49, wherein the 38 amino acids in the second trefoil domain of the homologue amino acid sequence are identical to amino acids 58-95 of SEQ ID NO:1.

51. (New) The polypeptide of claim 50, wherein the 39 amino acids in the first trefoil domain of the homologue amino acid sequence are identical to amino acids 8-46 of SEQ ID NO:1.
52. (New) The polypeptide of claim 49, wherein the 39 amino acids in the first trefoil domain of the homologue amino acid sequence are identical to amino acids 8-46 of SEQ ID NO:1.
53. (New) The polypeptide of claim 42, wherein the homologue amino acid sequence is identical to SEQ ID NO:1 except for two amino acid substitutions.
54. (New) The polypeptide of claim 53, wherein the two amino acid substitutions are in the first trefoil domain of the homologue amino acid sequence.
55. (New) The polypeptide of claim 53, wherein the two amino acid substitutions are in the second trefoil domain of the homologue amino acid sequence.
56. (New) The polypeptide of claim 42, wherein the homologue amino acid sequence is identical to SEQ ID NO:1 except for deletion of one or more amino acids at either end of SEQ ID NO:1.
- E2 57. (New) The polypeptide of claim 56, wherein the homologue amino acid sequence is identical to SEQ ID NO:1 except for deletion of one amino acid at either end of SEQ ID NO:1.
58. (New) The polypeptide of claim 42, wherein the homologue amino acid sequence is identical to SEQ ID NO:1 except for addition of one or more amino acids at either end of SEQ ID NO:1.
59. (New) The polypeptide of claim 58, wherein the homologue amino acid sequence is identical to SEQ ID NO:1 except for addition of one amino acid at either end of SEQ ID NO:1.
60. (New) The polypeptide of claim 40, wherein said polypeptide is glycosylated at an Asn present at position 15 of the amino acid sequence of said polypeptide.
61. (New) The polypeptide of claim 60, wherein the glycosylated form comprises a glycosylated side chain comprising at least one hexose unit.
62. (New) The polypeptide of claim 61, wherein the glycosylated side chain comprises at least one mannose unit.
63. (New) The polypeptide of claim 62, wherein the glycosylated side chain comprises 13-17 mannose units.
64. (New) The polypeptide of claim 63, wherein the glycosylated form comprises at least one unit of N-acetyl glucosamine (GlcNAc).
65. (New) The polypeptide of claim 64, wherein the glycosylated form comprises (GlcNAc)<sub>2</sub>(Man)<sub>10-15</sub>.